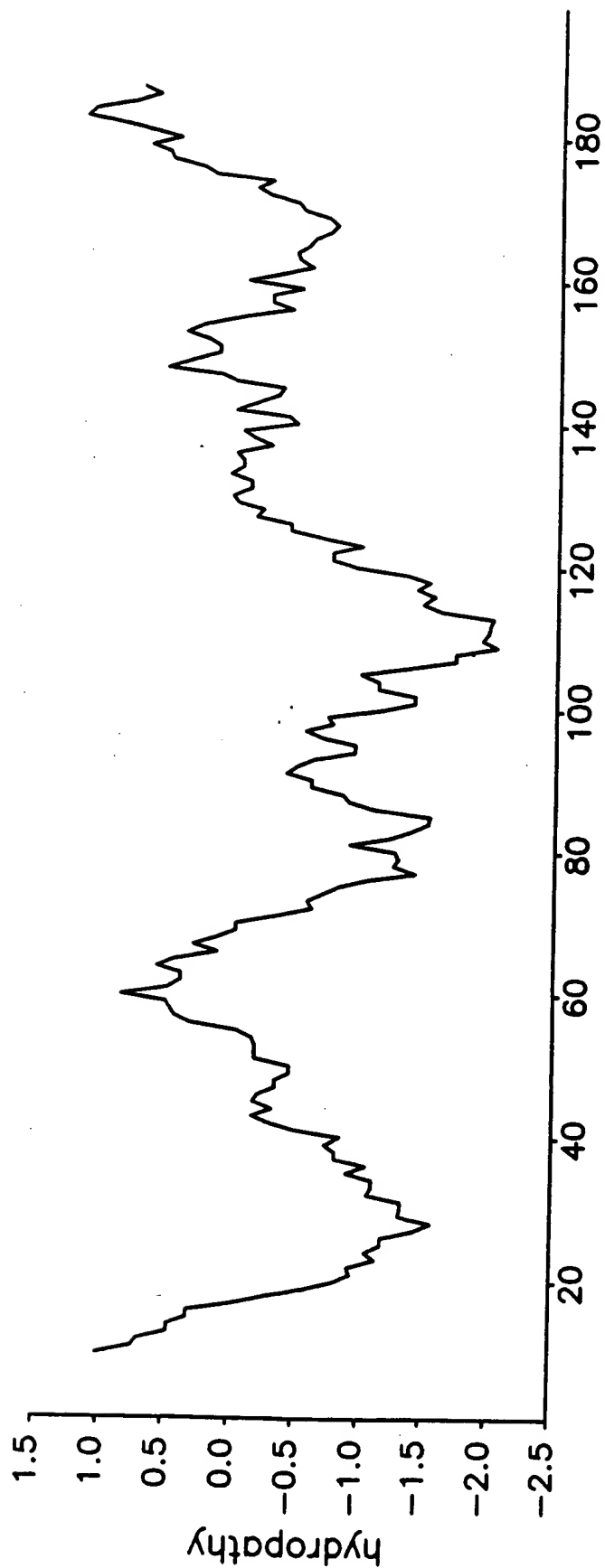


home/ruby/va/Molbio/nico/DNA6/p1.DNA62377 (length: 197)  
kyte (hydropathy); window: 20



IL 17 biology #2

DNA 62377

```
> /nico Unix Files/DNA6/ss.DNA62377
> 0 Sites [No Sites]
> length: 1047 bp (circular)
```

```
> /usr/seqdb2/sst/DNA/Dnaseqs.full/ss.DNA62377 (1250 bases)
[DNA62377], sheldens
```

>insert starts here

```
1 GCCAGGTGTG CAGGCCGCTC CAAGCCACGC CTGCCCCGGT GCCGCCACCA TGACGCTCCT CCGCGCCTC CTGTTTCTGA CCTGGCTGCA CACATGCCTG
  CGGTCCACAC GTCCGGCGAG GTTCGGGTG GACGGGCGGA CGCGGTGGT ACTGCGAGGA GGGCGGAG GACAAAGACT GGACCGACGT GTGTACGGAC
1
      M T L L P G L L F L T W L H T C L
      ^MET

101 GCCCACCATG ACCCTCCCT CAGGGGGCAC CCCCACAGTC ACGGTACCC ACACGTCTAC TCGGCTGAGG AACTGCCCT CGGCCAGGCC CCCCCACACC
    CGGTGTGTAC TGGGGAGGA GTCCCCCGTG GGGGTGTCTG TGCCATGGG TGTCACGATG AGCCGACTCC TTGACGGGGA GCCGGTCCGG GGGGGTGTGG
18 A H H D P S L R G H P H S H G T P H C Y S A E E L P L G Q A P P H L

201 TGCTGGCTCG AGGTGCCAAG TGGGGGCAGG CTTTGCCTGT AGCCTGGTG TCCAGCCTGG AGGCAGCAAG CCACAGGGGG AGGCACGAGA GGCCCTCAGC
    ACGACCGAGC TCCACGGTTC ACCCCGTCC GAAACGGACA TCGGGACCAC AGGTGGAGC TCCGTCTGTC GGTGTCCCC TCCGTGCTCT CCGGGAGTCG
52 L A R G A K W G Q A L P V A L V S S L E A A S H R G R H E R P S A

301 TAGCACCCAG TGCCCGGTGC TGCGGCCGGA GGAGGTGTTG GAGGCAGACA CCACACAGCG CTCCATCTCA CCCTGGAGAT ACCGTGTGGA CACGGATGAG
    ATGCTGGTTC ACGGGCCACG ACGCCGGCCT CCTCCACAAC CTCGGTCTGT GGGTGGTGGC GAGGTAGAT GGGACCTCTA TGGCACACCT GTGCCTACTC
85 T T Q C P V L R P E E V L E A D T H Q R S I S P W R Y R V D T D E

401 GACCGCTATC CACAGAAGCT GGCCTTCGCC GAGTGCCTGT GCAGAGGCTG TATCGATGCA CGGACGGGCC GCGAGACAGC TGCCTCAAC TCCGTGCGGC
    CTGGCGATAG GTGTCTTCGA CCGGAAGCGG CTCACGGACA CGTCTCCGAC ATAGCTACGT GCCTGCCCGG CGCTCTGTG ACGCAGATTG AGGCACGCCG
118 D R Y P Q K L A F A E C L C R G C I D A R T G R E T A A L N S V R L

501 TGCTCCAGAG CTGTGTTGTTG CTGCGCCGCC GGCCCTGCTC CCGGACGGC TCGGGGCTCC CCACACCTGG GGCCTTTGCC TTCCACACCG AGTTTCATCCA
    ACGAGGTCTC GGACGACCAC GACGCGCGG GCGGACGAG GCGGTGCGG AGCCCGAGG GGTGTGGACC CCGGAAACGG AAGGTGTGGC TCAAGTAGGT
152 L Q S L L V L R R R P C S R D G S G L P T P G A F A F H T E F I H

601 CGTCCCCGTC GGCTGCACCT GCGTGTGTC CCGTTTCAGT TGACCGCGGA GGGCGTGGG CCCCTAGACT GGACACGTGT GCTCCCCAGA GGGCACCCCC
    GCAGGGGCAG CCGACGTGGA CGCACGACGG GGCAAGTCAC ACTGGCGGCT CCGGCACCCC GGGGATCTGA CCTGTGCACA CGAGGGGTCT CCCGTGGGGG
185 V P V G C T C V L P R S V O

701 TATTATGTG TATTATTGT TATTATATG CCTCCCCCAA CACTACCTT GGGTCTGGG CATTCCCCGT GTCTGGAGGA CAGCCCCCA CTGTTCTCCT
    ATAAATACAC ATAAATAACA ATAAATATAC GGAGGGGGTT GTGATGGAA CCCCAGACCC GTAAGGGCA CAGACCTCCT GTCGGGGGT GACAAGAGA
```

801 CATCTCCAGC CTCAGTAGTT GGGGGTAGAA GGAGCTCAGC ACCTCTTCCA GCCCTTAAAG CTGCAGAAA GGTGTCACAC GGCTGCCTGT ACCTTGGCTC  
 GTAGAGGTCG GAGTCATCAA CCCCATCTT CCTCGAGTCG TGGAGAAGGT CCGGAATTTC GACGCTTTT CCACAGTGT CCGACGGACA TGGAAACCGAG  
 901 CCTGTCTCTGC TCCCGGCTTC CCTTACCCTA TCACTGGCCT CAGGCCCCGC AGGCTGCCTC TTCCCAACCT CCTTGGAAAGT ACCCCTGTTT CTTAAACAAT  
 GGACAGGACG AGGCCGAAG GGAATGGAT AGTGACCGGA GTCCGGGGCG TCCGACGGAG AAGGGTTGGA GGAACCTTCA TGGGGACAAA GAATTTGTTA  
 1001 TATTTAAGTG TACGTGTATT ATTAACTGA TGAACACATC CCCAAA  
 ATAAATTCAC ATGCACATAA TAATTGACT ACTTGTGTAG GGGTTTT